RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:

Source:

Date Processed by STIC:

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial	Number: 10 578 580 CRF Edit Date: 5-22- Edited by: 72
	Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line
	Corrected the SEQ ID NO. Sequence numbers edited were:
	Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
<u>V</u>	Deleted:invalid beginning/end-of-file text ; page numbers
	Inserted mandatory headings/numeric identifiers, specifically:
	Moved responses to same line as heading/numeric identifier, specifically:
	Other:

Revised 09/09/2003



IFWP

RAW SEQUENCE LISTING DATE: 05/22/2006
PATENT APPLICATION: US/10/578,580 TIME: 09:47:22

Input Set : A:\PTO.KD.txt

```
3 <110> APPLICANT: AMGEN INC.
             Aldrich, Teri
      5
              Shen, Wenyan
      6
              Jacobsen, Frederick W.
      7
              Morris, Arvia E.
      8
              Allen, Martin J.
     10 <120> TITLE OF INVENTION: Monkey Immunoglobulin Sequences
     12 <130> FILE REFERENCE: A-951 (WO)
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/578,580
C--> 15 <141> CURRENT FILING DATE: 2006-05-05
     17 <150> PRIOR APPLICATION NUMBER: US 60/517,970
     18 <151> PRIOR FILING DATE: 2003-11-07
     20 <160> NUMBER OF SEQ ID NOS: 86
     22 <170> SOFTWARE: PatentIn version 3.2
     24 <210> SEQ ID NO: 1
     25 <211> LENGTH: 999
     26 <212> TYPE: DNA
     27 <213> ORGANISM: Macaca fascicularis
     29 <400> SEQUENCE: 1
     30 geetecaeca agggeeeate ggtetteece etggegeeet eetecaggag caceteegag
                                                                               60
     32 ageacagegg ceetgggetg cetggteaag gactaettee etgaaceegt gacegtgteg
                                                                              120
     34 tggaactcag gctccctgac cageggcgtg cacaccttcc cggctgtcct acagtcctca
                                                                              180
     36 gggetetaet eceteageag egtggtgaee gtgeeeteea geagettggg caeceagaee
                                                                              240
                                                                              300
     38 tacgtctgca acgtaaacca caagcccagc aacaccaagg tggacaagag agttgagata
     40 aaaacatgtg gtggtggcag caaacctccc acgtgcccac cgtgcccagc acctgaactc
                                                                              360
     42 ctggggggac cgtcagtctt cctcttcccc ccaaaaccca aggacaccct catgatetcc
                                                                              420
     44 cggacccctg aggtcacgtg cgtggtggta gacgtgagcc aggaagaccc cgatgtcaag
                                                                              480
     46 ttcaactggt acgtaaatgg cgcggaggtg catcatgccc agacgaagcc acgggagacg
                                                                              540
     48 cagtacaaca gcacatateg tgtggtcage gteetcaeeg teaegeaeca ggaetggetg
                                                                              600
     50 aacggcaagg agtacacgtg caaggtctcc aacaaagccc tcccggcccc catccagaaa
                                                                              660
     52 accateteca aagacaaagg geageeeega gageeteagg tgtacaeeet geeeeegtee
                                                                              720
     54 cgggaggagc tgaccaagaa ccaggtcagc ctgacctgcc tggtcaaagg cttctacccc
                                                                              780
     56 agegacateg tegtggagtg ggagageage gggcageegg agaacaceta caagaceace
                                                                              840
     58 cegecegtge tggaeteega eggeteetae tteetetaea geaageteae egtggaeaag
                                                                              900
                                                                              960
     60 agcaggtggc agcaggggaa cgtcttctca tgctccgtga tgcatgaggc tctgcacaac
     62 cactacaccc agaagagcct ctccctgtct ccgggtaaa
                                                                              999
     65 <210> SEQ ID NO: 2
     66 <211> LENGTH: 333
     67 <212> TYPE: PRT
     68 <213> ORGANISM: Macaca fascicularis
     70 <400> SEQUENCE: 2
     72 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Pro Ser Ser Arg
     73 1
                        5
                                            10
```

RAW SEQUENCE LISTING DATE: 05/22/2006
PATENT APPLICATION: US/10/578,580 TIME: 09:47:22

Input Set : A:\PTO.KD.txt

```
76 Ser Thr Ser Glu Ser Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
                                   25
80 Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ser Leu Thr Ser
                               40
84 Gly Val His Thr Phe Pro Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser
                           55
88 Leu Ser Ser Val Val Thr Val Pro Ser Ser Leu Gly Thr Gln Thr
                       70
                                            75
92 Tyr Val Cys Asn Val Asn His Lys Pro Ser Asn Thr Lys Val Asp Lys
96 Arg Val Glu Ile Lys Thr Cys Gly Gly Ser Lys Pro Pro Thr Cys
97
                                   105
               100
100 Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser Val Phe Leu
            115
                                120
104 Phe Pro Pro Lys Pro Lys Asp Thr Leu Met Ile Ser Arg Thr Pro Glu
                            135
                                                 140 ·
108 Val Thr Cys Val Val Val Asp Val Ser Gln Glu Asp Pro Asp Val Lys
                        150
                                            155
112 Phe Asn Trp Tyr Val Asn Gly Ala Glu Val His His Ala Gln Thr Lys
                    165
                                        170
116 Pro Arg Glu Thr Gln Tyr Asn Ser Thr Tyr Arg Val Val Ser Val Leu
                180
                                    185
120 Thr Val Thr His Gln Asp Trp Leu Asn Gly Lys Glu Tyr Thr Cys Lys
            195
                                200
124 Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Gln Lys Thr Ile Ser Lys
                            215
125
                                                 220
128 Asp Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu Pro Pro Ser
                        230
                                            235
132 Arg Glu Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys Leu Val Lys
                    245
                                        250
136 Gly Phe Tyr Pro Ser Asp Ile Val Val Glu Trp Glu Ser Ser Gly Gln
                260
                                    265
140 Pro Glu Asn Thr Tyr Lys Thr Thr Pro Pro Val Leu Asp Ser Asp Gly
            275
                                280
144 Ser Tyr Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser Arg Trp Gln
                            295
148 Gln Gly Asn Val Phe Ser Cys Ser Val Met His Glu Ala Leu His Asn
                                            315
                                                                 320
                        310
152 His Tyr Thr Gln Lys Ser Leu Ser Leu Ser Pro Gly Lys
153
                    325
156 <210> SEQ ID NO: 3
157 <211> LENGTH: 1581
158 <212> TYPE: DNA
159 <213> ORGANISM: Macaca fascicularis
161 <400> SEQUENCE: 3
162 gcctccacca agggcccatc ggtcttcccc ctggcgtcct gctccaggag cacctcccag
                                                                           60
164 agcacagegg cectgggetg cetggteaag gactaettee eegaaceegt gacegtgteg
                                                                          120
166 tggaactcag gcgccctgac cagcggcgtg cacaccttcc aggctgtcct acagtcctca
                                                                          180
168 gggetetaet eecteageag egtggtgaee gtgeeeteea geagettggg eacteagaee
                                                                          240
```

RAW SEQUENCE LISTING DATE: 05/22/2006
PATENT APPLICATION: US/10/578,580 TIME: 09:47:22

Input Set : A:\PTO.KD.txt

```
170 tacgtctgca acgtcgttca tgagcccagc aacaccaagg tggacaagac agttggtgag
                                                                          300
172 aggccagcga gggaaggggg gtgtctgctg gaagccaggc tcggccctcc tgcctggaca
                                                                          360
174 aactetgget gtgeageece ageecaggge ageagggeag geecegtetg tetteteace
                                                                          420
176 cagaggeete tgeecaccee acteatgete agggageeag tettetgget ttttecacea
                                                                          480
178 ggetetgage aggeacagge tggatgeece taccecagge cetgeacaca caggggeagg
                                                                          540
                                                                          600
180 tgctgggctc agacctgcca agagccatat ctgggaggac cctgccctga cctaagccca
182 ccccaaaggc caaactccac tccctcagct cagacacctt ctctcctccc acatcccagt
                                                                          660
184 aacteceaat ettetetetg eagggeteee atgtegttee aegtgeeeac egtgeeeagg
                                                                          720
186 taagccagec caggeeteac cetecagete aaggtgggac aagegeeeta gagtggeetg
                                                                          780
188 tgtccaggga caggccctgc ccgggtgctg acacgtccac ctccatctct tcctcagctg
                                                                          840
190 aactcctggg gggaccgtca gtcttcctct tccccccaaa acccaaggac accctcatga
                                                                          900
192 tttcccggac ccctgaggtc acgtgcgtgg tggtagacgt gagccaggaa gaacccgatg
                                                                          960
194 tcaagttcaa ctggtacgtg gacggcgtgg aggtgcacaa tgcccagacg aagccacggg
                                                                         1020
196 aggagcagtt caacagcacg taccgcgtgg tcagcgtcct caccgtcaca caccaggact
                                                                         1080
198 ggctgaacgg caaggagtac acgtgcaagg tctccaacaa agccctcccg gccccaaagc
                                                                         1140
200 agaaaactgt ctccaaaacc aaaggtggga cccgcggggc acgagggcca cgtggacaga
                                                                         1200
202 ggccggctca gcccaccctc tgccctggga gtgaccgctg tgccaacctc tgtccctaca
                                                                         1260
204 gggcagcccc gagagccaca ggtgtacacc ctgcccccgc cccgggagga gctgaccaag
                                                                         1320
206 aaccaggtca gcctgacctg cctggtcaaa ggcttctacc ccagcgacat cgtcgtggag
                                                                         1380
208 tgggcgagca acgggcagcc ggagaacacc tacaagacca ccccgcccgt gctggactcc
                                                                         1440
210 gacggctcct acttcctcta cagcaagetc accgtggaca agagcaggtg gcagcagggg
                                                                         1500
212 aacacettet catgeteegt gatgeatgag getetgeaca accaetacae ceagaagage
214 ctctccgtgt ctccgggtaa a
                                                                         1581
217 <210> SEQ ID NO: 4
218 <211> LENGTH: 326
219 <212> TYPE: PRT
220 <213> ORGANISM: Macaca fascicularis
222 <400> SEQUENCE: 4
224 Ala Ser Thr Lys Gly Pro Ser Val Phe Pro Leu Ala Ser Cys Ser Arg
228 Ser Thr Ser Gln Ser Thr Ala Ala Leu Gly Cys Leu Val Lys Asp Tyr
229
                20
                                    25
232 Phe Pro Glu Pro Val Thr Val Ser Trp Asn Ser Gly Ala Leu Thr Ser
           35
                                40
236 Gly Val His Thr Phe Gln Ala Val Leu Gln Ser Ser Gly Leu Tyr Ser
       50
                            55
240 Leu Ser Ser Val Val Thr Val Pro Ser Ser Leu Gly Thr Gln Thr
                                            75
245 Tyr Val Cys Asn Val Val His Glu Pro Ser Asn Thr Lys Val Asp Lys
246
                                        90
249 Thr Val Gly Leu Pro Cys Arg Ser Thr Cys Pro Pro Cys Pro Ala Glu
250
                100
                                    105
253 Leu Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp
                                120
257 Thr Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Asp
       130
                            135
                                                140
261 Val Ser Gln Glu Glu Pro Asp Val Lys Phe Asn Trp Tyr Val Asp Gly
                        150
265 Val Glu Val His Asn Ala Gln Thr Lys Pro Arg Glu Glu Gln Phe Asn
```

RAW SEQUENCE LISTING DATE: 05/22/2006
PATENT APPLICATION: US/10/578,580 TIME: 09:47:22

Input Set : A:\PTO.KD.txt

```
266
                    165
                                         170
269 Ser Thr Tyr Arg Val Val Ser Val Leu Thr Val Thr His Gln Asp Trp
                180
                                     185
273 Leu Asn Gly Lys Glu Tyr Thr Cys Lys Val Ser Asn Lys Ala Leu Pro
            195
                                 200
                                                     205
277 Ala Pro Lys Gln Lys Thr Val Ser Lys Thr Lys Gly Gln Pro Arg Glu
                            215
                                                 220
281 Pro Gln Val Tyr Thr Leu Pro Pro Pro Arg Glu Glu Leu Thr Lys Asn
                        230
                                             235 -
285 Gln Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile
                    245
                                         250
289 Val Val Glu Trp Ala Ser Asn Gly Gln Pro Glu Asn Thr Tyr Lys Thr
290
                260
                                     265
293 Thr Pro Pro Val Leu Asp Ser Asp Gly Ser Tyr Phe Leu Tyr Ser Lys
                                 280
297 Leu Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Thr Phe Ser Cys
        290
                            295
                                                 300
301 Ser Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu
                        310
                                             315
306 Ser Val Ser Pro Gly Lys
307
                    325
310 <210> SEQ ID NO: 5
311 <211> LENGTH: 1579
312 <212> TYPE: DNA
313 <213> ORGANISM: Macaca fascicularis
315 <400> SEQUENCE: 5
316 geetecaeca agggeecate ggtetteece etggegteet eetecaggag eaceteegag
                                                                            60
318 agcacagegg ccctgggctg cctggtcaag gactacttcc ccgaacccgt gactgtgtcg
                                                                          120
320 tggaactcag gcgccctgac cagcggcgtg cacaccttcc cggctgtcct acagtcctca
                                                                          180
322 gggetetaet eceteageag egtggtgaee gtgeeeteea geagettggg eaceeagaee
                                                                          240
324 tacgtctgca acgtcgttca tgagcccagc aacaccaagg tggacaagag agttggtgag
                                                                          300
326 aggccagega gggagggga gtgtctgctg gaagccatgc tcggccctcc tgcctggaca
                                                                          360
328 aaccetgget gtgeageece ageceaggge ageagggeag geeeggtetg teteeteace
                                                                          420
330 cagaggeete tgeecaceee aeteatgete agggagaeag tettetgget tttteeaeea
                                                                          480
332 gactccgagc aggcacaggc tggatgcccc taccccaggc tctgcacaca taggggctgg
                                                                          540
334 tgctgggctc agacctgcca agagccatat ctgggaggac cctgctcctg acctaagccc
                                                                          600
336 accccaaagg ccaaactcca ctccctcagc tcggaaacct tctctcctac cagatcccag
                                                                          660
338 taactcccaa tettetetet geagagttea cacccccatg cecaccatge ceaggtaage
                                                                          720
340 cagcccaggc ctcgccctcc agctcaaggt gggacaagtg ccctagagtg gcctgtgtcc
                                                                          780
342 agggacaggc cccgcctggg tgctgacatg cccacctcca tctcttcctc agcacctgaa
                                                                          840
344 ctcctggggg gaccgtcagt cttcctcttc cccccaaaac ccaaqgacac cctcatgatc
                                                                          900
346 tcccggaccc ctgaggtcac atgcgtggtg gtggacgtga gccaggaaga ccccgaggtc
                                                                          960
348 cagttcaact ggtacgtgga cggcgtggag gtgcatcatg cccagacgaa gccacgggag
                                                                         1020
350 aggcagttca acagcacgta ccgcgtggtc agcgtcctca ccgtcacaca ccaggactgg
                                                                         1080
352 ctgaacggca aggagtacac gtgcaaggtc tccaacaaag gcctcccggc ccccatcgag
                                                                         1140
354 aaaaccatct ccaaagccaa aggtgggacc cgcggggccc gagggccacg tggacagagg
                                                                         1200
356 ccggctcagc ccaccctctg ccctgggagt gaccgctgtg ccaacctctg tccctacagg
                                                                         1260
358 gcagccccga gagccgcagg tgtacatect gcccccgccc caggaggagc tgaccaagaa
                                                                         1320
360 ccaggtcage ctgacctgcc tggtcacagg cttctacccc agegacateg ccgtggagtg
                                                                         1380
```

RAW SEQUENCE LISTING DATE: 05/22/2006 PATENT APPLICATION: US/10/578,580 TIME: 09:47:23

Input Set : A:\PTO.KD.txt

364 366 368 371 372	ggagagcaac gggcagccgg agaacaccta caagaccacc ccgcccgtgc tggactccga cggctcctac ttcctctaca gcaagctcat cgtggacaag agcaggtggc agcaggggaa caccttctca tgctccgtga tgcatgaggc tctgcacaac cactacaccc agaagagcct ctccctgtct ccgggtaaa <210> SEQ ID NO: 6 <211> LENGTH: 325 <212> TYPE: PRT														1440 1500 1560 1579		
374	<21	3 > O	RGAN:	ISM:	Maca	aca :	fasc:	icula	aris								
376	<40	0 > S	EQUE	NCE:	6												
		Ser	Thr	Lys	Gly	Pro	Ser	Val	Phe	Pro	Leu	Ala	Ser	Ser	Ser	Arg	
379			_		5				_	10	_	_		_	15	_	
	Ser	Thr	Ser		Ser	Thr	Ala	Ala		Gly	Cys	Leu	Val	_	Asp	Tyr	
383	D1	D	a 1	20	*** 7	m\	77-7	a	25		0	~1	31 -	30	m1	a	
	Pne	Pro		Pro	vaı	Thr	vaı	Ser	Trp	Asn	Ser	GIY		ьeu	Thr	ser	
387	C111	77-1	35	Пр×	Dho	Dro	. ה ר ת	40	T 011	~1 _n	Cox	C 0 x	45	T 011	Пт гж	Cor	
390	GIY	50	птэ	1111	PIIE	PIO	55	Val	Leu	GIII	ser	60	GIY	ьец	ıyı	ser	
	T.=11		Ser	Val	Val	Thr		Pro	Sar	Ser	Sar		Gl v	Thr	Gln	Thr	
395		OCI	DCI	Vai	Val	70	val	110	DCI	SCI	75	шец	Gry	1111	GIII	80	
		Val	Cvs	Asn	Val		His	Glu	Pro	Ser		Thr	Lvs	Val	Asp		
399	-1-		-1-		85					90			-1-		95	-2-	
	Arq	Val	Glu	Phe	Thr	Pro	Pro	Cys	Pro		Cys	Pro	Ala	Pro		Leu	•
403	_			100				•	105		•			110			
406	Leu	Gly	Gly	Pro	Ser	Val	Phe	Leu	Phe	Pro	Pro	Lys	Pro	Lys	Asp	Thr	
407			115					120					125				
410	Leu	Met	Ile	Ser	Arg	Thr	Pro	Glu	Val	Thr	Cys	Val	Val	Val	Asp	Val	
411		130					135					140					
414	Ser	Gln	Glu	Asp	Pro	Glu	Val	Gln	Phe	Asn	\mathtt{Trp}	Tyr	Val	Asp	Gly		
	145					150					155					160	
	Glu	Val	His	His		Gln	Thr	Lys	Pro	_	Glu	Arg	Gln	Phe		Ser	
419	-1				165	_		_	_,	170	1		~~	_	175	_	
	Thr	ıyr	Arg		vaı	Ser	vaı	Leu		vaı	Thr	His	Gin	_	Trp	Leu	
423	7 00	C111	Tva	180	Пэ гэс	Πh~	Ctra	T	185	Cox	7.00	T	c1	190	Dro	71-	
427	ASII	GIY	195	GIU	ıyı	1111	Cys	Lys 200	vai	ser	ASII	гу	205	ьеи	PIO	Ala	
	Pro	Tle		Lvs	Thr	Tle	Ser	Lys	Δla	Taye	Glv	Gln		Δra	Glu	Pro	
431	110	210	GIU	цуБ	1111	110	215	цуз	лта	цуз	Gry	220	110	AL 9	GIU	110	
	Gln		Tvr	Ile	Leu	Pro		Pro	Gln	Glu	Glu		Thr	Lvs	Asn	Gln	
	225		-1-			230					235			-1-		240	
		Ser	Leu	Thr	Cys		Val	Thr	Gly	Phe		Pro	Ser	Asp	Ile		
439					245				-	250	-			-	255		
442	Val	Glu	Trp	Glu	Ser	Asn	Gly	Gln	Pro	Glu	Asn	Thr	Tyr	Lys	Thr	Thr	
443				260					265					270			
446	${\tt Pro}$	Pro		Leu	Asp	Ser	Asp	Gly	Ser	${\tt Tyr}$	Phe	Leu	Tyr	Ser	Lys	Leu	
447			275					280					285				
	Ile		Asp	Lys	Ser	Arg	_	Gln	Gln	Gly	Asn		Phe	Ser	Cys	Ser	
451	_	290	•			_	295	_		_		300	_	_	_	_	
		Met	His	Glu	Ala		His	Asn	His	Tyr		Gln	Lys	Ser	Leu		
455	305					310					315					320	

VERIFICATION SUMMARY

DATE: 05/22/2006

PATENT APPLICATION: US/10/578,580

TIME: 09:47:24

Input Set : A:\PTO.KD.txt

Output Set: N:\CRF4\05222006\J578580.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application Number L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date

Raw Sequence Listing before editing (for reference only)



IFWP

RAW SEQUENCE LISTING DATE: 05/17/2006
PATENT APPLICATION: US/10/578,580 TIME: 10:15:14

Input Set: A:\A-951 (WO) SeqListing final 10.19.04.txt
Output Set: N:\CRF4\05172006\J578580.raw

```
3 <110> APPLICANT: AMGEN INC.
      4
            Aldrich, Teri
      5
              Shen, Wenyan
              Jacobsen, Frederick W.
      6
      7
             Morris, Arvia E.
      8
             Allen, Martin J.
     10 <120> TITLE OF INVENTION: Monkey Immunoglobulin Sequences
     12 <130> FILE REFERENCE: A-951 (WO)
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/578,580
C--> 15 <141> CURRENT FILING DATE: 2006-05-05
     17 <150> PRIOR APPLICATION NUMBER: US 60/517,970
     18 <151> PRIOR FILING DATE: 2003-11-07
     20 <160> NUMBER OF SEQ ID NOS: 86
```

22 <170> SOFTWARE: PatentIn version 3.2

Does Not Comply
Corrected Diskette Needed-(pq, 1)

ERRORED SEQUENCES

```
3483 <210> SEQ ID NO: 86
     3484 <211> LENGTH: 109
     3485 <212> TYPE: PRT
     3486 <213> ORGANISM: Artificial Sequence
     3488 <220> FEATURE:
     3489 <223> OTHER INFORMATION: Antibody variable domain sequences that recognize anti IL-
4R
     3491 <400> SEQUENCE: 86
     3493 Asp Ile Val Leu Thr Gln Thr Pro Ala Thr Leu Ser Leu Ser Pro Gly
                                              10
     3497 Glu Arg Ala Thr Leu Ser Cys Arg Ala Ser Gln Ser Val Gly Ser Ser
     3498
                      20
                                          25
     3501 Tyr Leu Ala Trp Tyr Gln Gln Arg Pro Gly Gln Ala Pro Arg Leu Leu
     3505 Ile Tyr Gly Ala Ser Ser Arg Ala Thr Gly Ile Pro Asp Arg Phe Ser
     3509 Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Arg Leu Glu
     3510 65
     3513 Pro Glu Asp Phe Ala Val Tyr Tyr Cys Gln Gln Tyr Gly Ser Ser Pro
     3514
                                              90
     3517 Pro Trp Met Phe Gly Gln Gly Thr Lys Val Glu Ile Lys
     3518
                      100
E--> 3521/A-951 (WO)
E--> 3529 Express Mail Label No. EV 531746269 US deleted
E--> 3526 37
```

VERIFICATION SUMMARY

DATE: 05/17/2006

PATENT APPLICATION: US/10/578,580

TIME: 10:15:16

Input Set : A:\A-951 (WO) SeqListing final 10.19.04.txt

Output Set: N:\CRF4\05172006\J578580.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application Number

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:3521 M:333 E: Wrong sequence grouping, Amino acids not in groups!

L:3521 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:2

L:3526 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:86

L:3529 M:333 E: Wrong sequence grouping, Amino acids not in groups!

L:3529 M:330 E: (2) Invalid Amino Acid Designator, NUMBER OF INVALID KEYS:6

L:3529 M:252 E: No. of Seq. differs, <211> LENGTH:Input:109 Found:117 SEQ:86 /